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ERROR REPORT

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Application Serial Number: _09/424,686_

Source: _1652_

Date Processed by STIC: _2/14/2002_

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
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    U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
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Revised 01/29/2002

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1652

RAW SEQUENCE LISTING   DATE: 02/14/2002
PATENT APPLICATION: US/09/424,686   TIME: 15:18:45

Input Set : A:\pto.vsk.txt
Output Set: N:\CRF3\02142002\I424686.raw Does Not Comply
                                        Corrected Diskette Needed

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    4 <110> APPLICANT: Hagen, Gustav
    5         Siegmund, Hans-Ulrich
    6         Weichel, Walter
    7         Wick, Maresa
    8         Zubov, Dmitry
   10 <120> TITLE OF INVENTION: Human Catalytic Telomerase Sub-Unit and its Diagnostic and
Therapeutic
   11         Use
   13 <130> FILE REFERENCE: Bayer 10,203
   15 <140> CURRENT APPLICATION NUMBER: US 09/424,686
   17 <141> CURRENT FILING DATE: 1999-11-29
   19 <150> PRIOR APPLICATION NUMBER: PCT/EP98/03468
   21 <151> PRIOR FILING DATE: 1998-06-09
   23 <160> NUMBER OF SEQ ID NOS: 7
   25 <170> SOFTWARE: Microsoft Word
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ERRORED SEQUENCES

Invalid response must be PRT for proteins.

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  171 <210> SEQ ID NO: 2
  172 <211> LENGTH: 1132
E--> 173 <212> TYPE: amino acid
  174 <213> ORGANISM: Homo sapiens
  176 <400> SEQUENCE: 2
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  178     1               5                   10                  15
  180     His Tyr Arg Glu Val Leu Pro Leu Ala Thr Phe Val Arg Arg Leu Gly
  181                 20                  25                  30
  183     Pro Gln Gly Trp Arg Leu Val Gln Arg Gly Asp Pro Ala Ala Phe Arg
  184                 35                  40                  45
  186     Ala Leu Val Ala Gln Cys Leu Val Cys Val Pro Trp Asp Ala Arg Pro
  187             50                  55                  60
  189     Pro Pro Ala Ala Pro Ser Phe Arg Gln Val Ser Cys Leu Lys Glu Leu
  190     65                  70                  75                  80
  192     Val Ala Arg Val Leu Gln Arg Leu Cys Glu Arg Gly Ala Lys Asn Val
  193                 85                  90                  95
  195     Leu Ala Phe Gly Phe Ala Leu Leu Asp Gly Ala Arg Gly Gly Pro Pro
  196                 100                 105                 110
  198     Glu Ala Phe Thr Thr Ser Val Arg Ser Tyr Leu Pro Asn Thr Val Thr
  199                 115                 120                 125
  201     Asp Ala Leu Arg Gly Ser Gly Ala Trp Gly Leu Leu Leu Arg Arg Val
  202             130                 135                 140
  204     Gly Asp Asp Val Leu Val His Leu Leu Ala Arg Cys Ala Leu Phe Val
  205     145                 150                 155                 160
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PATENT APPLICATION: US/09/424,686   TIME: 15:18:45

Input Set : A:\pto.vsk.txt
Output Set: N:\CRF3\02142002\I424686.raw

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208                 165                 170                 175
210     Gln Leu Gly Ala Ala Thr Gln Ala Arg Pro Pro Pro His Ala Ser Gly
211                 180                 185                 190
213     Pro Arg Arg Arg Leu Gly Cys Glu Arg Ala Trp Asn His Ser Val Arg
214                 195                 200                 205
216     Glu Ala Gly Val Pro Leu Gly Leu Pro Ala Pro Gly Ala Arg Arg Arg
217                 210                 215                 220
219     Gly Gly Ser Ala Ser Arg Ser Leu Pro Leu Pro Lys Arg Pro Arg Arg
220     225                 230                 235                 240
222     Gly Ala Ala Pro Glu Pro Glu Arg Thr Pro Val Gly Gln Gly Ser Trp
223                 245                 250                 255
225     Ala His Pro Gly Arg Thr Arg Gly Pro Ser Asp Arg Gly Phe Cys Val
226                 260                 265                 270
228     Val Ser Pro Ala Arg Pro Ala Glu Glu Ala Thr Ser Leu Glu Gly Ala
229                 275                 280                 285
231     Leu Ser Gly Thr Arg His Ser His Pro Ser Val Gly Arg Gln His His
232                 290                 295                 300
234     Ala Gly Pro Pro Ser Thr Ser Arg Pro Pro Arg Pro Trp Asp Thr Pro
235     305                 310                 315                 320
237     Cys Pro Pro Val Tyr Ala Glu Thr Lys His Phe Leu Tyr Ser Ser Gly
238                 325                 330                 335
240     Asp Lys Glu Gln Leu Arg Pro Ser Phe Leu Leu Ser Ser Leu Arg Pro
241                 340                 345                 350
243     Ser Leu Thr Gly Ala Arg Arg Leu Val Glu Thr Ile Phe Leu Gly Ser
244                 355                 360                 365
246     Arg Pro Trp Met Pro Gly Thr Pro Arg Arg Leu Pro Arg Leu Pro Gln
247         370                 375                 380
249     Arg Tyr Trp Gln Met Arg Pro Leu Phe Leu Glu Leu Leu Gly Asn His
250     385                 390                 395                 400
252     Ala Gln Cys Pro Tyr Gly Val Leu Leu Lys Thr His Cys Pro Leu Arg
253                 405                 410                 415
255     Ala Ala Val Thr Pro Ala Ala Gly Val Cys Ala Arg Glu Lys Pro Gln
256                 420                 425                 430
258     Gly Ser Val Ala Ala Pro Glu Glu Glu Asp Thr Asp Pro Arg Arg Leu
259             435                 440                 445
261     Val Gln Leu Leu Arg Gln His Ser Ser Pro Trp Gln Val Tyr Gly Phe
262         450                 455                 460
264     Val Arg Ala Cys Leu Arg Arg Leu Val Pro Pro Gly Leu Trp Gly Ser
265     465                 470                 475                 480
267     Arg His Asn Glu Arg Arg Phe Leu Arg Asn Thr Lys Lys Phe Ile Ser
268                 485                 490                 495
270     Leu Gly Lys His Ala Lys Leu Ser Leu Gln Glu Leu Thr Trp Lys Met
271                 500                 505                 510
273     Ser Val Arg Asp Cys Ala Trp Leu Arg Arg Ser Pro Gly Val Gly Cys
274                 515                 520                 525
276     Val Pro Ala Ala Glu His Arg Leu Arg Glu Glu Ile Leu Ala Lys Phe
277         530                 535                 540
279     Leu His Trp Leu Met Ser Val Tyr Val Val Glu Leu Leu Arg Ser Phe
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280    545                    550                    555                    560
282    Phe Tyr Val Thr Glu Thr Thr Phe Gln Lys Asn Arg Leu Phe Phe Tyr
283                    565                    570                    575
285    Arg Lys Ser Val Trp Ser Lys Leu Gln Ser Ile Gly Ile Arg Gln His
286                    580                    585                    590
288    Leu Lys Arg Val Gln Leu Arg Glu Leu Ser Glu Ala Glu Val Arg Gln
289                    595                    600                    605
291    His Arg Glu Ala Arg Pro Ala Leu Leu Thr Ser Arg Leu Arg Phe Ile
292                    610                    615                    620
294    Pro Lys Pro Asp Gly Leu Arg Pro Ile Val Asn Met Asp Tyr Val Val
295    625                    630                    635                    640
297    Gly Ala Arg Thr Phe Arg Arg Glu Lys Arg Ala Glu Arg Leu Thr Ser
298                  . 645                    650                    655
300    Arg Val Lys Ala Leu Phe Ser Val Leu Asn Tyr Glu Arg Ala Arg Arg
301                    660                    665                    670
303    Pro Gly Leu Leu Gly Ala Ser Val Leu Gly Leu Asp Asp Ile His Arg
304                    675                    680                    685
306    Ala Trp Arg Thr Phe Val Leu Arg Val Arg Ala Gln Asp Pro Pro Pro
307                    690                    695                    700
309    Glu Leu Tyr Phe Val Lys Val Asp Val Thr Gly Ala Tyr Asp Thr Ile
310    705                    710                    715                    720
312    Pro Gln Asp Arg Leu Thr Glu Val Ile Ala Ser Ile Ile Lys Pro Gln
313                    725                    730                    735
315    Asn Thr Tyr Cys Val Arg Arg Tyr Ala Val Val Gln Lys Ala Ala His
316                    740                    745                    750
318    Gly His Val Arg Lys Ala Phe Lys Ser His Val Ser Thr Leu Thr Asp
319                    755                    760                    765
321    Leu Gln Pro Tyr Met Arg Gln Phe Val Ala His Leu Gln Glu Thr Ser  /
322                    770                    775                    780
324    Pro Leu Arg Asp Ala Val Val Ile Glu Gln Ser Ser Ser Leu Asn Glu
325    785                    790                    795                    800
327    Ala Ser Ser Gly Leu Phe Asp Val Phe Leu Arg Phe Met Cys His His
328                    805                    810                    815
330    Ala Val Arg Ile Arg Gly Lys Ser Tyr Val Gln Cys Gln Gly Ile Pro
331                    820                    825                    830
333    Gln Gly Ser Ile Leu Ser Thr Leu Leu Cys Ser Leu Cys Tyr Gly Asp
334                    835  .                  840                    845
336    Met Glu Asn Lys Leu Phe Ala Gly Ile Arg Arg Asp Gly Leu Leu Leu
337                    850                    855                    860
339    Arg Leu Val Asp Asp Phe Leu Leu Val Thr Pro His Leu Thr His Ala
340    865                    870                    875                    880
342    Lys Thr Phe Leu Arg Thr Leu Val Arg Gly Val Pro Glu Tyr Gly Cys
343                    885                    890                    895
345    Val Val Asn Leu Arg Lys Thr Val Val Asn Phe Pro Val Glu Asp Glu
346                    900                    905                    910
348    Ala Leu Gly Gly Thr Ala Phe Val Gln Met Pro Ala His Gly Leu Phe
349                    915                    920                    925
351    Pro Trp Cys Gly Leu Leu Leu Asp Thr Arg Thr Leu Glu Val Gln Ser
352                    930                    935                    940
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RAW SEQUENCE LISTING   DATE: 02/14/2002
PATENT APPLICATION: US/09/424,686   TIME: 15:18:45

Input Set : A:\pto.vsk.txt
Output Set: N:\CRF3\02142002\I424686.raw

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354    Asp Tyr Ser Ser Tyr Ala Arg Thr Ser Ile Arg Ala Ser Leu Thr Phe
355    945             950             955             960
357    Asn Arg Gly Phe Lys Ala Gly Arg Asn Met Arg Arg Lys Leu Phe Gly
358                965             970             975
360    Val Leu Arg Leu Lys Cys His Ser Leu Phe Leu Asp Leu Gln Val Asn
361            980             985             990
363    Ser Leu Gln Thr Val Cys Thr Asn Ile Tyr Lys Ile Leu Leu Leu Gln
364         995             1000            1005
366    Ala Tyr Arg Phe His Ala Cys Val Leu Gln Leu Pro Phe His Gln Gln
367         1010            1015            1020
369    Val Trp Lys Asn Pro Thr Phe Phe Leu Arg Val Ile Ser Asp Thr Ala
370    1025            1030            1035            1040
372    Ser Leu Cys Tyr Ser Ile Leu Lys Ala Lys Asn Ala Gly Met Ser Leu
373                1045            1050            1055
375    Gly Ala Lys Gly Ala Ala Gly Pro Leu Pro Ser Glu Ala Val Gln Trp
376                1060            1065            1070
378    Leu Cys His Gln Ala Phe Leu Leu Lys Leu Thr Arg His Arg Val Thr
379            1075            1080            1085
381    Tyr Val Pro Leu Leu Gly Ser Leu Arg Thr Ala Gln Thr Gln Leu Ser
382         1090            1095            1100
384    Arg Lys Leu Pro Gly Thr Thr Leu Thr Ala Leu Glu Ala Ala Ala Asn
385    1105            1110            1115            1120
387    Pro Ala Leu Pro Ser Asp Phe Lys Thr Ile Leu Asp
388                1125            1130
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VERIFICATION SUMMARY
PATENT APPLICATION: US/09/424,686

DATE: 02/14/2002
TIME: 15:18:46

Input Set : **A:\pto.vsk.txt**
Output Set: **N:\CRF3\02142002\I424686.raw**

L:173 M:310 E: (3) Wrong or Missing Sequence Type, TYPE: